

seq list.txt  
SEQUENCE LISTING

&lt;110&gt; The Ontario Cancer Institute

&lt;120&gt; Cytotoxic Heteromeric Protein Combinatorial Libraries

&lt;130&gt; 34104-0079

&lt;140&gt; 09/601,644

&lt;141&gt; 1998-12-08

&lt;150&gt; PCT/CA98/01137

&lt;151&gt; 1998-12-08

&lt;150&gt; CA2,222,993

&lt;151&gt; 1998-02-04

&lt;160&gt; 4

&lt;170&gt; PatentIn Ver. 2.1

&lt;210&gt; 1

&lt;211&gt; 293

&lt;212&gt; PRT

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; Description of Artificial Sequence:Primer

&lt;400&gt; 1

Lys Glu Phe Thr Leu Asp Phe Ser Thr Ala Lys Thr Tyr Val Asp Ser  
1 5 10 15Leu Asn Val Ile Arg Ser Ala Ile Gly Thr Pro Leu Gln Thr Ile Ser  
20 25 30Ser Gly Gly Thr Ser Leu Leu Met Ile Asp Ser Gly Ser Gly Asp Asn  
35 40 45Leu Phe Ala Val Asp Val Arg Gly Ile Asp Pro Glu Glu Gly Arg Phe  
50 55 60Asn Asn Leu Arg Leu Ile Val Glu Arg Asn Asn Leu Tyr Val Thr Gly  
65 70 75 80Phe Val Asn Arg Thr Asn Asn Val Phe Tyr Arg Phe Ala Asp Phe Ser  
85 90 95

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His	Val	Thr	Phe	Pro	Gly	Thr	Thr	Ala	Val	Thr	Leu	Ser	Gly	Asp	Ser		
			100					105					110				
Ser	Tyr	Thr	Thr	Leu	Gln	Arg	Val	Ala	Gly	Ile	Ser	Arg	Thr	Gly	Met		
		115					120					125					
Gln	Ile	Asn	Arg	His	Ser	Leu	Thr	Thr	Ser	Tyr	Leu	Asp	Leu	Met	Ser		
	130					135					140						
His	Ser	Gly	Thr	Ser	Leu	Thr	Gln	Ser	Val	Ala	Arg	Ala	Met	Leu	Arg		
145					150					155					160		
Phe	Val	Thr	Val	Thr	Ala	Glu	Ala	Leu	Arg	Phe	Arg	Gln	Ile	Gln	Arg		
				165					170					175			
Gly	Phe	Arg	Thr	Thr	Leu	Asp	Asp	Leu	Ser	Gly	Arg	Ser	Tyr	Val	Met		
			180					185					190				
Thr	Ala	Glu	Asp	Val	Asp	Leu	Thr	Leu	Asn	Trp	Gly	Arg	Leu	Ser	Ser		
		195					200					205					
Val	Leu	Pro	Asp	Tyr	His	Gly	Gln	Asp	Ser	Val	Arg	Val	Gly	Arg	Ile		
	210					215					220						
Ser	Phe	Gly	Ser	Ile	Asn	Ala	Ile	Leu	Gly	Ser	Val	Ala	Leu	Ile	Leu		
225					230					235					240		
Asn	Cys	His	His	His	Ala	Ser	Arg	Val	Ala	Arg	Met	Ala	Ser	Asp	Glu		
				245					250					255			
Phe	Pro	Ser	Met	Cys	Pro	Ala	Asp	Gly	Arg	Val	Arg	Gly	Ile	Thr	His		
			260					265					270				
Asn	Lys	Ile	Leu	Trp	Asp	Ser	Ser	Thr	Leu	Gly	Ala	Ile	Leu	Met	Arg		
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Arg	Thr	Ile	Ser	Ser													
		290															

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<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:Primer

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Thr Pro Asp Cys Val Thr Gly Lys Val Glu Tyr Thr Lys Tyr Asn Asp  
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Asp Asp Thr Phe Thr Val Lys Val Gly Asp Lys Glu Leu Phe Thr Asn  
20 25 30

Arg Trp Asn Leu Gln Ser Leu Leu Leu Ser Ala Gln Ile Thr Gly Met  
35 40 45

Thr Val Thr Ile Lys Thr Asn Ala Cys His Asn Gly Gly Gly Phe Ser  
50 55 60

Glu Val Ile Phe Arg  
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<211> 94

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:Primer

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<210> 4

<211> 128

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:Primer

<400> 4

tacgtactgc agctcgagtc aacgaaaaat aacttcgctg aatccaccgc cattatggca 60  
cgcgtaggtt ttaatgggta cagtcatacc ggtaatttgc gcactgagaa gaagagactg 120  
aagattcc 128

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